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		THE STREET PROJECTION

(54) Title: ADENOVIRAL VECTORS INCLUDING DNA ENCODING LUNG SURFACTANT PROTEIN

An adenoviral vector including a DNA sequence encoding a lung surfactant protein. The adenoviral vector may be a replication deficient adenoviral vector which is free of at least the majority of the E1 and E3 DNA sequences. Such vectors may be employed for generation of infectious viral particles which may transduce lung epithelial cells in vivo to enable the expression of lung surfactant protein by such cells.

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ADENOVIRAL VECTORS INCLUDING DNA ENCODING LUNG SURFACTANT PROTEIN

This application is a continuation-in-part of application Serial No. 08/044,406, filed April 8, 1993.

This invention relates to adenoviral vectors. More particularly, this invention relates to adenoviral vectors which include DNA encoding a lung surfactant protein and to the use of such vectors in treating disease states associated with lung surfactant protein deficiency, such as infant respiratory distress syndrome, and adult respiratory distress syndrome.

Surfactant proteins are natural endogenous proteins produced primarily within the alveolar and airway epithelial cells of the normal lung and interact with phospholipids to maintain the patency of the alveolar structures. When pulmonary surfactant protein concentration on the alveolar surface falls below critical levels, surface tension of the liquid-gas interface increases, thereby leading to alveolar collapse, pulmonary ventilation-perfusion mismatch, and hypoxia. In severe cases, this can lead to death. Intermittent administration of exogenous bovine lung surfactant protein has shown partial, but not

complete remission of the pathophysiology of the surfactant deficiency state.

It is an object of the present invention to provide a recombinant expression vehicle for expressing pulmonary surfactant protein.

It is a further object of the present invention to provide an expression vehicle which will enable prolonged expression of pulmonary surfactant protein in the lung in order to correct the clinical surfactant protein deficiency state and its attendant pathophysiologic effects on gas exchange.

The above objects and others should be apparent from the following specification.

In accordance with an aspect of the present invention, there is provided an adenoviral vector including a DNA sequence encoding a lung surfactant protein.

In one embodiment, the adenoviral vector is a replication deficient adenoviral vector, i.e., such vector is free of a DNA sequence(s) which is (are) required for viral replication, such as, for example, the El DNA sequence or a portion thereof. In one embodiment, the adenoviral vector is free of at least a portion of the adenoviral El DNA sequence and is free of at least a portion of the adenoviral El DNA sequence. The E3 region encodes several polypeptides which help the adenovirus to evade the immune surveillance of the host.

In one embodiment, the adenoviral vector comprises an adenoviral 5' inverted terminal repeat, or ITR; an adenoviral 3'

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encoding a lung surfactant protein; and a promoter controlling the expression of the DNA sequence encoding the lung surfactant protein. The vector is free of at least the majority of the adenoviral E1 and E3 DNA sequences, but is not free of all of the E2 and E4 DNA sequences, and is not free of DNA sequences encoding adenoviral proteins expressed by the adenoviral major late promoter. In one embodiment, the vector is also free of at least a portion of at least one DNA sequences selected from the group consisting of the E2 and E4 DNA sequences. In another embodiment, the vector is free of the adenoviral E1 and E3 DNA sequences, and is free of one of the E2 and E4 DNA sequences, and is free of one of the E2 and E4 DNA sequences, and is free of one of the E2 and E4 DNA sequences.

In yet another embodiment, the vector is free of at least the majority of the El and E3 DNA sequences, is free of at least a portion of at least one DNA sequence selected from the group consisting of the E2 and E4 DNA sequences, and is free of DNA sequences encoding adenoviral proteins expressed under control of the adenoviral major late promoter.

The DNA sequence encoding a lung surfactant protein is under the control of a suitable promoter. Suitable promoters which may be employed include, but are not limited to, adenoviral promoters, such as the adenoviral major late promoter; or heterologous promoters, such as the cytomegalovirus (CMV) promoter; the Rous Sarcoma Virus promoter; the respiratory

syncytial virus promoter; inducible promoters, such as the mouse mammary tumor virus, or MMTV, promoter; the metallothionein promoter; and heat shock promoters. In addition, tissue-specific promoters such as, but not limited to, lung surfactant protein promoters, may also be employed. It is to be understood, however, that the scope of the present invention is not to be limited to any specific promoter.

Lung surfactant proteins which may be encoded by the DNA sequence encoding a lung surfactant protein include surfactant protein A (SPA), surfactant protein B (SPB), and surfactant protein C (SPC).

Surfactant protein A is described in Kuroki, et al., J.

Biol. Chem., Vol. 263, No. 7, pgs. 3388-3394 (March 5, 1988).

Surfactant protein B and DNA encoding therefor are described in Pilot-Matias, et al., DNA, Vol. 8, No. 2, pgs. 75-86 (1989),

Glasser, et al., Proc. Nat. Acad. Sci., Vol 84, pgs. 4007-4011 (June 1987); Revak, et al., J. Clin. Invest., Vol. 81, pgs. 826-833 (March 1988); O'Reilly, et al., Biochimica et Biophysica Acta, Vol. 1011, pgs. 140-148 (1989); and Weaver, et al., J.

Amer. Phys. Soc., pgs. 982-987 (1988). Surfactant protein C is described further in Glasser, et al., J. Biol. Chem., Vol. 263, No. 21, pgs. 10326-10331 (July 25, 1988).

In one embodiment, the DNA sequence encoding a lung surfactant protein encodes lung surfactant protein B. Present evidence suggests that SPB is the most clinically important lung surfactant protein of those herein above described.

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Such a vector, in a preferred embodiment, is assembled first by constructing, according to standard techniques, a shuttle plasmid which contains, beginning at the leftward adenoviral genomic elements, the "critical left end elements", which include an adenoviral 5' ITR, an adenoviral encapsidation signal, and the Ela enhancer sequence; a promoter (which may be an adenoviral promoter or a foreign promoter); a tripartite leader sequence, a multiple cloning site; a poly A signal; and a DNA segment which corresponds to a segment of the adenoviral genome. Such DNA segment serves as a substrate for homologous recombination with a modified or mutated adenovirus, and such sequence may encompass, for example, a segment of the adenoviral genome from base 3328 to base 6241 of the adenovirus 5 genome. The plasmid may also include a selectable marker and an origin of replication. The origin of replication, may be, for example, a bacterial origin of replication. A representative example of such a shuttle plasmid is pAVS6, shown in Figure 4. An intron may be included within the transcribed portion to enhance the cytoplasmic mRNA accumulation levels.

The multiple cloning site facilitates the insertion of the DNA sequence encoding a lung surfactant protein into the plasmid. The DNA sequence encoding the lung surfactant protein may be inserted into the multiple cloning site. In general, restriction enzyme sites separating the above-mentioned components of the shuttle plasmid include "rare" restriction enzyme sites; i.e., sites which are found to occur randomly in

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eukaryotic genes at a frequency from about one in every 10,000 to about one in every 100,000 base pairs. This increases the flexibility and ease of rearranging components of the vectors in assembled shuttle plasmids.

Homologous recombination is then effected with a modified or mutated adenovirus in which at least the majority of the El and E3 adenoviral DNA sequences have been deleted, as shown, for example, in Figure 8. Such homologous recombination may be effected through co-transfection of the shuttle plasmid and the modified adenovirus into a helper cell line, such as 293 (embryonic kidney epithelial) cells, by CaPO4 precipitation. Upon such homologous recombination, a cloning vector is formed in which the modified adenovirus DNA which was 5' to the DNA segment in the shuttle plasmid corresponding to a similar segment of the modified adenoviral genome is replaced with the components in the shuttle plasmid which are 5' to such DNA segment. This homologous recombination, or "crossing over" event, can occur anywhere along the segment of the genome of the modified adenovirus which corresponds to the segment which is also contained within the shuttle plasmid (such as, for example, bases 3328 to 6241 of adenovirus 5 in Example 1 shown below).

Through such homologous recombination, a vector is formed which includes an adenoviral 5' ITR; an adenoviral encapsidation signal; an Ela enhancer sequence; a promoter; a tripartite leader sequence; a DNA sequence encoding a lung surfactant protein; a poly A signal; adenoviral DNA free of at least the majority of

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the El and E3 adenoviral DNA sequences; and an adenoviral 3' ITR.

This vector may then be introduced into a cell line such as the

293 cell line for production of large amounts of infectious

recombinant adenoviral particles. The 293 cell line is a human

fetal kidney epithelial cell line into which has been permanently

introduced 11% of the left end of the adenovirus 5 genome. This

directs the synthesis of the adenoviral E1a and E1b proteins and

allows trans-complementation of E1-deleted vectors.

The infectious viral particles may then be administered to a host in vivo as part of a gene therapy procedure. Such infectious viral particles may be administered systemically, such as by intravenous or intraperitoneal or intrasmuscular or subcutaneous administration, or may be administered topically, such as by intratracheal or intrabronchial administration, or, alternatively, the infectious viral particles may be administered in an aerosol formulation. The infectious viral particles may be administered in an amount of up to about 10¹³ pfu, preferably from about 10⁷ pfu to about 10¹² pfu. For example, the infectious viral particles may be employed in the transduction of the epithelium of the respiratory tract or alveoli; whereby the lung epithelial cells will express lung surfactant protein in amounts sufficient to achieve clinical correction of lung surfactant protein deficiency.

In addition, the infectious viral particles may be used to transduce eukaryotic cells in vitro. Eukaryotic cells which may be transduced include, but are not limited to, macrophages,

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lymphocytes, fibroblasts, liver cells, bronchial cells, and other epithelial or endothelial cells. Such eukaryotic cells then may be administered to a host as part of a gene therapy procedure, or may be cultured in vitro whereby such cells produce lung surfactant protein.

In addition, the infectious viral particles may be used to transduce eukaryotic cells in vitro for the in vitro production of lung surfactant protein. Examples of eukaryotic cells which may be transduced in vitro for the in vitro production of lung surfactant protein include, but are not limited to, those eukaryotic cells hereinabove described, as well as Chinese Hamster Ovary (CHO) cells, COS-7 cells, NIH 3T3 cells, vero cells, HeLa cells, MRC-5 cells, CN1 cells, W138 cells, and chicken lymphoma cells. The lung surfactant protein produced by such cells may then be administered to a host in conjunction with an acceptable pharmaceutical carrier in order to treat lung surfactant protein deficiency states.

In another embodiment, the vector comprises an adenoviral 5' ITR; an adenoviral 3' ITR; an adenoviral encapsidation signal; a DNA sequence encoding a lung surfactant protein; and a promoter controlling the DNA sequence encoding a lung surfactant protein. The vector is free of the adenoviral E1, E2, E3, and E4 DNA sequences, and the vector is free of DNA sequences encoding adenoviral proteins promoted by the adenoviral major late promoter; i.e., the vector is free of DNA encoding adenoviral

structual proteins. Such vector is sometimes hereinafter referred to as a "gutless adenoviral vector," or "GLAd" vector.

Promoters which are contained in the vector may be those hereinabove described.

Such vectors may be constructed by removing the adenoviral 5' ITR, the adenoviral 3' ITR, and the adenoviral encapsidation signal, from an adenoviral genome by standard techniques. Such components, as well as a promoter (which may be an adenoviral promoter or a non-adenoviral promoter), tripartite leader sequence, poly A signal, may, by standard techniques, be ligated into a base plasmid or "starter" plasmid such as, for example, pKSII (Strategene), to form an appropriate cloning vector. The cloning vector may include a multiple cloning site, as hereinabove described, to facilitate the insertion of the foreign DNA sequence into the cloning vector. An appropriate vector in accordance with the present invention is thus formed by cutting the cloning vector by standard techniques at appropriate restriction sites in the multiple cloning site, and then ligating the DNA sequence encoding a lung surfactant protein into the cloning vector.

The GLAd vector may then be packaged into infectious viral particles using a helper adenovirus or cell line which provides the necessary packaging materials. If a helper virus is used, in one embodiment, preferably it has a defective encapsidation signal in order that the helper virus will not package itself. Examples of such encapsidation-defective helper viruses which may

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be employed are described in Grable, et al., <u>J. Virol.</u>, Vol. 66, pgs. 723-731 (1992), and in Grable, et al., <u>J. Virol.</u>, Vol. 64, pgs. 2047-2056 (1990). In another embodiment, the helper virus has a normal packaging signal.

DNA for the vector and the encapsidation-defective helper virus are transfected into an appropriate cell line for the generation of infectious viral particles. Transfection may take place by electroporation, calcium phosphate precipitation, microinjection, or through proteoliposomes. Examples of appropriate cell lines include, but are not limited to, HeLa cells, A549 cells, or 293 (embryonic kidney epithelial) cells. The infectious viral particles may then be purified away from helper virus by CsCl isopycnic density centrifugation and transduced into lung epithelial cells lining the respiratory tract or alveoli, as hereinabove described, whereby such cells express lung surfactant protein.

In another alternative, the vector is transfected into the cells, followed by infection of the cells with the encapsidation-defective helper virus.

The invention will now be described with respect to the following example, however, the scope of the present invention is not intended to be limited thereby.

Example 1

The adenoviral construction shuttle plasmid pAvS6 was constructed in several steps using standard cloning techniques including polymerase chain reaction based cloning techniques.

First, the 2913 bp BglII, HindIII fragment was removed from Addis27 and inserted as a blunt fragment into the XhoI site of pKSII (Stratagene, La Jolla, CA) (Figure 1). Ad-d1327 (Thimmappaya, et al., Cell, Vol. 31, pg. 543 (1983)) is identical to adenovirus 5 except that an XbaI fragment including bases 28593 to 30470 (or map units 78.5 to 84.7) of the adenovirus 5 genome, and which is located in the E3 region, has been deleted.

The comlete Adenovirus 5 genome is registered as Genbank accession #M73260, incorporated herein by reference, and the virus is available from the American Type Culture Collection, Rockville, Maryland, U.S.A. under accession number VR-5.

Ad-d1327 was constructed by routine methods from Adenovirus 5 (Ad5). The method is outlined briefly as follows and previously described by Jones and Shenk, Cell 13:181-188 (1978). Ad5 DNA is isolated by proteolytic digestion of the virion and partially cleaved with Xba I restriction endonuclease. The Xba I fragments are then reassembled by ligation as a mixture of fragments. This results in some ligated genomes with a sequence similar to Ad5, except excluding sequences 28593 bp to 30470 bp. This DNA is then transfected into suitable cells (e.g. KB cells, HeLa cells, 293 cells) and overlaid with soft agar to allow plaque formation. Individual plaques are then isolated, amplified, and screened for the absence of the 1878 bp E3 region Xba 1 fragment. The orientation of this fragment was such that the BgIII site was nearest the T7 RNA polymerase site of pKSII

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and the HindIII site was nearest the T3 RNA polymerase site of pKSII. This plasmid was designated pHR. (Figure 1).

Second, the ITR, encapsidation signal, Rous Sarcoma Virus promoter, the adenoviral tripartite leader (TPL) sequence and linking sequences were assembled as a block using PCR amplification (Figure 2). The ITR and encapsidation signal (sequences 1-392 of Ad-d1327 [identical to sequences from Ad5, Genbank accession #M73260]) were amplified (amplification 1) together from Ad-d1327 using primers containing NotI or AscI restriction sites. The Rous Sarcoma Virus LTR promoter was amplified (amplification 2) from the plasmid pRC/RSV (sequences 209 to 605; Invitrogen, San Diego, CA) using primers containing an AscI site and an SfiI site. DNA products from amplifications 1 and 2 were joined using the "overlap" PCR method (amplification 3) with only the NotI primer and the SfiI primer. Complementarity between the AscI containing end of each initial DNA amplification product from reactions 1 and 2 allowed joining of these two pieces during amplification. Next the TPL was amplified (amplification 4) (sequences 6049 to 9730 of Ad-d1327 [identical to similar sequences from Ad5, Genbank accession #M73260]) from cDNA made from mRNA isolated from 293 cells infected for 16 hours with Ad-d1327 using primers containing Sfil and XbaI sites respectively. DNA fragments from amplification reactions 3 and 4 were then joined using PCR (amplification 5) with the NotI- and XbaI-site-containing primers, thus creating . the complete gene block.

Third, the ITR-encapsidation signal-TPL fragment was then purified, cleaved with NotI and XbaI and inserted into the NotI, XbaI cleaved pHR plasmid. This plasmid was designated pAvS6A and the orientation was such that the NotI site of the fragment was next to the T7 RNA polymerase site (Figure 3).

Fourth, the SV40 early polyA signal was removed from SV40 DNA as an HpaI-BamHI fragment, treated with T4 DNA polymerase and inserted into the SalI site of the plasmid pAvS6A-(Figure 3) to create pAvS6 (Figures 3 and 4).

a 2kb DNA fragment containing the whole human pulmonary surfactant protein B (SPB) cDNA (Figure 5) (Pilot-Matias, 1989) was obtained from plasmid pKC4-SPB (Figure 6) (Weaver, et al., J. Amer. Phys. Soc., pgs. L-95 to L-103 (1992)) by EcoRI digestion. This DNA fragment was isolated, purified, and then cloned into the EcoRV site of plasmid pAVS6. (Figure 4). Three identical clones with correct insertion of SPB cDNA were obtained. Such clones are named pAVS6-SPB#7, pAVS6-SPB#12, and pAVS6-SPB#13.

PAVS6-SPB#7 is shown in Figure 7. The orientation of the SPB DNA within the shuttle plasmid was obtained by evaluating the DNA sequences of the two termini of the SPB cDNA insert in the plasmid with primers derived from pAVS6.

The recombinant adenoviral vector AVISPB1 (Figure 8), containing SPB cDNA was constructed through homologous recombination between the Ad5 deletion mutant Ad-d1327 (Figure 8), and pAVS6 SPB#7. Homologous recombination, or "crossing over," occurs between Ad-d1327 and pAVS6.SPB#7, along the

segment common to both Ad-dl327 and pAVS6.SPB#7 which corresponds to bases 3328 to 6241 (or map units 9.24 to 17.34) of the adenovirus 5 genome. Ad-dl327 has a deleted E3 region in which base pairs 28593 to 30470 are absent (Thimmappaya, et al, Cell, Vol. 31, pgs. 543-551 (1982)). pAVS6SPB#7 contains an adenoviral 5' ITR, an origin of replication contained completely within the 5' ITR, an Ela enhancer and encapsidation signal, a Rous Sarcoma Virus promoter, an adenovirus 5 tripartite leader sequence and the 2kb human SPB cDNA including the entire protein coding sequence (nucleotides 1 to 1172), and the SV40 poly A signal.

Example 2

293 cells (ATCC No. CRL 1573) were infected with AVISPB1 at a multiplicity of infection (MOI) of 50 MOI units. At 12 hours post-infection, the cells were radiolabeled with ³⁵S-methionine (50μCi/ml) overnight. Identical amounts of labeled protein were used for immunoprecipitation with antisera against SPB.

Immunopreciptates were analyzed by SDS-polyacrylamide gel electrophoresis on 16% gel. The gels were fluorographed. C⁴⁴-labeled molecular weight markers and BioRad broad range molecular weight markers were used as size markers. As shown in Figure 9, lane 1 shows an uninfected control; lane 2 shows AVISPB1-infected 293 cells in which electrophoresis of immunoprecipitates occurred under reducing conditions; and lane 3 shows AVISPB1-infected 293 cells in which electrophoresis of immunoprecipitates occurred under non-reducing conditions. The active SPB peptide migrates at approximately Mr=6,000 to 8,000 (reduced) and forms oligomers

(unreduced) identical to that of native human SPB (Mr=18,000). The precursor protein also was detected in both reduced and unreduced conditions.

Example 3

Mouse lung Type II-like epithelial cell lines were transduced with adenoviral vector AVISPB1, which contains the full length human SPBcDNA under the control of the Rous Sarcoma Virus promoter. Expression of SPB was assessed by RNase protection using P-labelled probes specific for endogenous mouse SPB (upper band) or human SPB (middle band). A B-tubulin specific probe (lowest band) also was used to ensure that the same amount of RNA was added to each assay. The B-tubulin probe did not recognize human B-tubulin in lane 1. As shown in Figure 10, cells infected with AVISPB1 at 50 multiplicity of infection (MOI) units (lane 2), 100 MOI units (lane 3), and 150 MOI units (lane 4), clearly expressed human SPB mRNA. Human SPB was not detected in uninfected cells (lane 5).

Example 4

Four cotton rats were anesthetized by metaphane, and were given 1.5 x 10^{10} pfu of Av1SPB1. The rats were sacrificed at 2 (n=1), 3 (n=2) or 45 (n=1) days after administration, and the lungs were harvested. Total lung RNA then was extracted using the guanidine thiocyanate-CsCl technique (Chirgwin, et al., Biochemistry, Vol. 18, pgs. 5294-5299 (1979)). Total lung RNA(15 μ g) was subjected to formaldehyde agarose gel electrophoresis and transferred to a nylon membrane (Nytran,

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Schleicher and Schuell). The filter was crosslinked (UV Crosslink, Stratagene), and hybridized with a ³²P-labeled 2.0 kb human SP-B cDNA probe prepared by random priming (Loftstrand) and evaluated by autoradiography. Lungs from uninfected control rats and from rats infected with AvlLacZ4 also were subjected to the above hybridization procedure.

Northern blot analysis, as shown in Figure 11, demonstrated that human SP-B mRNA was expressed in the lungs of cotton rats infected with AvISP-B1, but not in those of uninfected animals nor in animals infected with AvILacZ4. A 2.0 kb mRNA for the human SP-B gene was detected in cotton rats infected with AvISP-B1, consistent with the size of the full length human SP-B mRNA transcribed by the vector.

Example 5

Two cotton rats were anesthetized with metaphane, and Av1SP-B1 (diluted in PBS to 10^{10} pfu/300 μ l) was administered via intranasal instillation. Two days after infection, the rats were sacrificed and the lungs were harvested. The lungs were washed twice in PBS and perfused with methionine-free LHC-8 medium (Biofluids), minced, and incubated for 24 hours in medium with 35 S-Cys/Met (1ml of medium, 100μ Ci/ml). The presence of SP-B in media and lysed lung explants was assessed by immunoprecipitation with rabbit anti-human SP-B antiserum to detect the secreted SP-B and processed SP-B peptides. (Weaver, et al., Am. J. Physiol., Vol. 263, pgs. L95-L103 (1992)). Av1Lacz4 treated rats and untreated rats were used as controls.



As shown in Figure 12, <u>de novo</u> synthesis and secretion of human SP-B peptide was detected from lung fragments removed from animals infected with Av1SPB1, and was not detected in uninfected animals or in animals infected with Av1LacZ4. Human SP-B peptide was detected as secreted, with 8 kda and 18 kda oligomeric forms suggesting that vector derived precursor SP-B (proSP-B) was processed after <u>in vivo</u> adenoviral vector-mediated gene transfer.

Example 6

Cotton rats were treated intranasally with AvISP-B1 in amounts of (a) 0.5 x 10° pfu (n=1); (b) 1.5 x 10° pfu (n=1); or (c) 1.5 x 10¹0 pfu (n=1). The animals were sacrificed 48 hours after infection, and the lungs were prepared for in situ hybridization analysis according to the method of Wert, et al., Development Biology, Vol. 156, pgs. 426-443 (1993), using human SP-BCRNA. An uninfected rat was used as a control.

Lung sections were inflation fixed in 20 cm of water in 4% paraformaldehyde at 4°C overnight. Radiolabeled ribroprobe for human SP-B was generated by in vitro transcription in the presence of 3°S-VTP. Slides were hybridized to the human SP-B probe under stringent conditions and exposed to emulsion from one to eight days, after which the slides were counterstained with hematoxylin and eosin. Antisense and sense probes were compared in Av1SPB1-treated, in Av1LacZ4-treated, and in control (untreated) animals.

In situ hybridization of the AvlSP-Bl infected cotton rats demonstrated expression of human SP-B mRNA in bronchiolar and

alveolar epithelial cells. Human SP-B mRNA was detected in a patchy distribution and increased in a dose-dependent manner, as shown in Figures 13 through 16, which show the in situ hybridization results of the rats treated with 0.5 x 10° pfu of Av1SP-B1; 1.5 x 10° pfu of Av1SP-B1; 1.5 x 10° pfu of Av1SP-B1; and an uninfected control rat, respectively. Light microscopic analysis of the lungs of Av1SP-B1 or Av1LacZ treated animals demonstrated a mild inflammatory response with a peribronchiclar lymphomonocytic infiltrate, increased macrophages, and some polymorpholeukocytes. The infiltrates were prominent after 48 hours and were dose-dependent. In parallel experiments with Av1lacZ4, the infiltrates were essentially resolved 3 to 4 weeks after exposure.

All patents, publications, and database entries referenced in this specification are indicative of the level of skill of persons in the art to which the invention pertains. The disclosures of all such patents, publications (including published patent applications), and database entries are specifically incorporated herein by reference in their entirety to the same extent as if each such individual patent, publication, and database entry were specifically and individually indicated to be incorporated by reference.

It is to be understood, however, that the scope of the present invention is not to be limited to the specific embodiments described above. The invention may be practiced

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other than as particularly described and still be within the scope of the accompanying claims.

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SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: Trapnell, Bruce

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(ii) TITLE OF INVENTION: Adenoviral Vectors Including

DNA Encoding Lung Surfactant

Protein

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- (V) COMPUTER READABLE FORM:
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 - (C) OPERATING SYSTEM: MS-DOS
 - (D) SOFTWARE: Word Perfect5.1

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 - (A) TELEPHONE: 201-994-1700
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- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 2,016 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: Genomic DNA

(ix) FEATURE:

(A) NAME/KEY: Surfactant protein B cDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

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					CCTTGGCCTG GGAACCGGAC	
					GCAGAGCCCT CGTCTCGGGA	
AGGGCATTGC	CTACAGGAAG	TCTGGGGACA	TGTGGGAGCC	GATGACCTAT	GCCAAGAGTG	240
TGAGGACATC	GTCCACATCC	TTAACAAGAT	GGCCAAGGAG	GCCATTTTCC	AGGACACGAT	300
	•				TCCTGTGCTA	
					ACGGGGTCAC ACCAGACTGA	
GTTGGTTCAC	GAACTGCTGA	TGAAGGGGGA	CCAGTAGCTG	ATGAAGGTCT	TGGTCTGACT	
					AGCCAGAGCA TCGGTCTCGT	
					CAGACCCTCT GTCTGGGAGA	
					CTGGGCCTCA GACCCGGAGT	
					GGCTCTGCAG CCGAGACGTC	
					TGGCAGTGGC ACCGTCACCG	
					TGGCTGAGCG ACCGACTCGC	
CTACTCCGTC	ATCCTGCTCG	ACACGCTGCT	GGGCCGCATG	CTGCCCCAGC	TGGTCTGCCG ACCAGACGGC	840
CCTCGTCCTC	CGGTGCTCCA	TGGATGACAG	CGCTGGCCCA	AGGTCGCCGA	CAGGAGAATG	900
					CCGGGAACAG	
CGACGGCGCT	CTGAGACTCA	CGGTGGAGAC	GTACAGGCAC	TGGTGGGTCC	GGCCCTTGTC GGCTGGACAG	•
					CCGACCTGTC	1020

GGAAAAGTGC AAGCAATTTG TGGAGCAGCA CACGCCCCAG CTGCTGACCC TGGTGCCCAG CCTTTTCACG TTCGTTAAAC ACCTCGTCGT GTGCGGGGTC GACGACTGGG ACCACGGGTC	1080
	1140
	1144
FEGUTGGGAT GCCCACACCA CCTGCCAGGC CCACGCCCAC ACACCCTGGT ACAGGTCGGG	
	1200
	1200
TCTCCAGTGT ATCCACAGCC CCGACCTTTG ATGAGAACTC AGCTGTCCAG CACGTTTCCT AGAGGTCACA TAGGTGTCGG GGCTGGAAAC TACTCTTGAG TCGACAGGTC CACGTTTCCT	
	1260
CAGAGGGGCT CTGGGACCAT GGTGACCAGG CICCICGGA CGAGGGACCG	
TTTCGGTTCA CTCTGGGGGG AAAAGAAGCC TCAGCTCCCA CACCGCCCTC CTCACCGCCCCCCCCCC	1320
TOCCACCTO AAAGAAGCC TCAGCTCCCA CACCGCCTC CACTGCGGG	
CCTCGCCAGC TGCCAGGCTG AAAAGAAGCC TCAGCTCCCA CACCGCCCTC GAGTGGCGGG GGAGCGGTCG ACGGTCCGAC TTTTCTTCGG AGTCGAGGGT GTGGCGGGAG GAGTGGCGGG GGAGCGGTCG ACGGTCCGAC TTTTCTTCGG AGTCGAGGGT GTGGCGGGAG GAGTGGCGGG	
GGAGCGGTCG ACCGTCCGAC TITTOTTCTC	1380
TTCCTCGGCA GTCACTTCCA CIGCACCTGG TGCCCGGGGG TCGGGACACAG GCCGGAACAG	·
	1440
TGTCTCAGCT CAACCACAGT CTGACACCAG AGCCCACTTC CATCCICIO COACACTCCG ACAGAGTCGA GTTGGTGTCA GACTGTGGTC TCGGGTGAAG GTAGGAGAGA CCACACTCCG ACAGAGTCGA GTTGGTGTCA GACTGTGGTC	
	1500
	1300
ACAGCGAGGG CAGCATCTGG AGGAGCTCTG CAGCCTCCAC ACCTACCACG	
	1560
GGAAAAACCA GCCACTGCTT TACAGGACAG GGGGTAGTTC GACTCGGGGC	
CCTCACACCC ACCCCATGC ACTCAAAGAT TGGATTTTAC AGCTACTTGC AATTCAAAAT CCTCACACCC ACCCCCATGC ACTCAAAGAT TGGATTTTAC TCGATGAACG TTAAGTTTTA	1620
ACCCCATGO ACTOAAAGAT TGGATTTTAC AGCTACTIOG ATTAGGTTTTA	
CCTCACACCC ACCCCCATGC ACTCAAAGAT TGGATTTTAC AGCTACTIGG TTAAGTTTTA GGAGTGTGGG TGGGGGTACG TGAGTTTCTA ACCTAAAATG TCGATGAACG TTAAGTTTTA	
GGAGTGTGGG TGGGGGTTGT CARACTGTTA	1680
GGAGTGTGGG TGGGGTATGGG AACATACAGA ACTCTAAAAG ATAGACATCA GAAATTGTTA TCAGAAGAAT AAAAAATGGG AACATACAGA ACTCTAAAAG ATAGACATCA CTTTAACAAT	
TCAGAAGAAT AAAAAATGGG AACATACAGA ACTCTAAAAG ATAGACATAT AGTCTTCTTA TTTTTTACCC TTGTATGTCT TGAGATTTTC TATCTGTAGT CTTTAACAAT	
AGTOTTOTTA TITILIACCO TITILIACCO	1740
AGTTAAGCTT TTTCAAAAAA ACTGGTTAAG GGTCGCATCA GTTCCCACCT GTACGTGCGCA	
	1800
TCTGGCATGA TGGGATGGCG ACCGGGCAAG CTTTCTTCCT GGAGATGGAG ACGACGAACT AGACCGTACT ACCCTACCGC TGGCCCGTTC GAAAGAAGGA GCTCTAGCAG ACGACGAACT	
	1860
GAGCTATTGC TTTGTTAAGA TATAAAAAGG GGTTTCTTTT TGTCTTTCTG AATTCCACCTG CTCGATAACG AAACAATTCT ATATTTTTCC CCAAAGAAAA ACAGAAAGAC ATTCCACCTGA	
	1920
	1320
TTCCAGCTTT TGATTGAAAG TCCTAGGGTG ATTCTATTTC TGCTGTGATT TATCTGCTGA TTCCAGCTTT TGATTGAAAG TCCTAGGGTG ATTCTATTTC TGCTGTGATT TATCTGCTGA AAGGTCGAAA ACTAACTTTC AGGATCCCAC TAAGATAAAG ACGACACTAA ATAGACGACC	
	1980
	1300
ANGETCAGET GGGGTTGTGC AAGCTAGGGA CCCATILCCAT ACATTATGTT ACAGACGTGG	
AAGCTCAGCT GGGGTTGTGC AAGCTAGGGA CCCATTCCTA TGTAATACAA ACAGACGTGG TTCGAGTCGA CCCCAACACG TTCGATCCCT GGGTAAGGAT ACATTATGTT ACAGACGTGG	7016
Titonatoe.	2016
AATGCTAATA AAGTCCTATT CTCTTTTATC GGAATT	
AATGCTAATA AAGTCCTAII CIAAAATAG CCTTAA TTACGATTAT TTCAGGATAA GAGAAAATAG CCTTAA	
TTACGALIAN	

WHAT IS CLAIMED IS:

- 1. An adenoviral vector including a DNA sequence encoding a lung surfactant protein.
- 2. The vector of Claim 1 wherein said lung surfactant protein is Surfactant Protein B.
- 3. The vector of Claim 1 wherein said adenoviral vector is a replication deficient adenoviral vector.
- 4. The vector of Claim 3 wherein said adenoviral vector is free of at least a portion of the adenoviral E1 DNA sequence, and is free of at least a portion of the adenoviral E3 DNA sequence.
- 5. The vector of Claim 4 wherein said vector comprises:
 - an adenoviral 5' ITR:
 - an adenoviral 3' ITR;
 - an adenoviral encapsidation signal;
 - a DNA sequence encoding a lung surfactant protein;
- a promoter controlling said foreign DNA sequence, said vector being free of at least the majority of the adenoviral E1 and E3 DNA sequences, but not free of all of the E2 and E4 DNA sequences and DNA sequences encoding adenoviral proteins controlled by the adenoviral major late promoter.
- 6. The vector of Claim 5 wherein the vector is free of at least the majority of the adenoviral E1 and E3 DNA sequences, and is free of at least a portion of at least one DNA sequence selected from the group consisting of the E2 and E4 DNA sequences.

7. The vector of Claim 6 wherein said vector is free of at least the majority of the adenoviral E1 and E3 DNA sequences, and is free of one of the E2 and E4 DNA sequences, and is free of a portion of the other of the E2 and E4 DNA sequences.

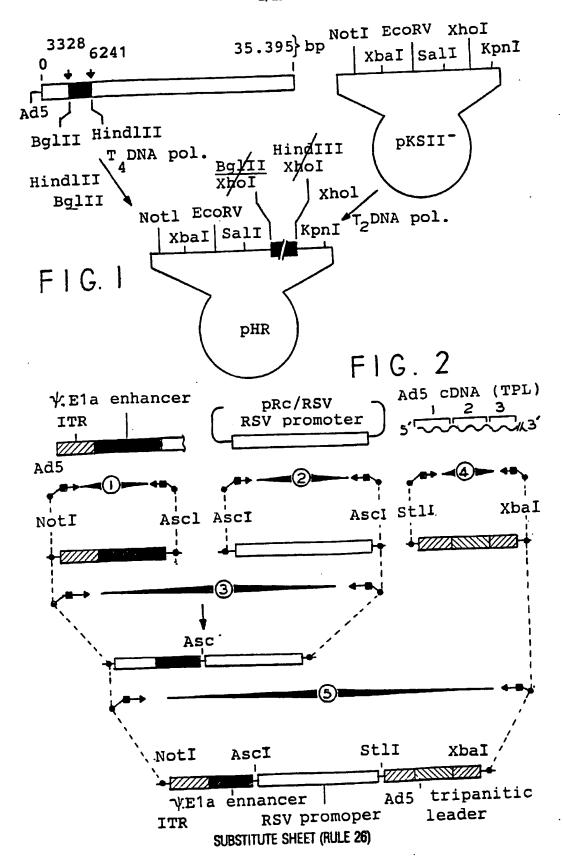
- 8. The vector of Claim 7 wherein said vector is free of at least the majority of the El and E3 DNA sequences, is free of at least a portion of at least one DNA sequence selected from the group consisting of the E2 and E4 DNA sequences, and is free of DNA sequences encoding adenoviral proteins promoted by the adenoviral major late promoter.
- 9. The vector of Claim 5 wherein said promoter is an adenoviral promoter.
- 10. The vector of Claim 5 wherein said promoter is a heterologous promoter.
- 11. Infectious viral particles generated from the vector of Claim 1.
- 12. Lung epithelial cells transfected with the infectious viral particles of Claim 11.
- 13. Infectious viral particles generated from the vector of Claim 5.
- 14. Lung epithelial cells transfected with the infectious viral particles of Claim 13.
- 15. A process for treating lung surfactant protein deficiency states, comprising:

administering to a host an effective amount of the infectious viral particles of Claim 11.

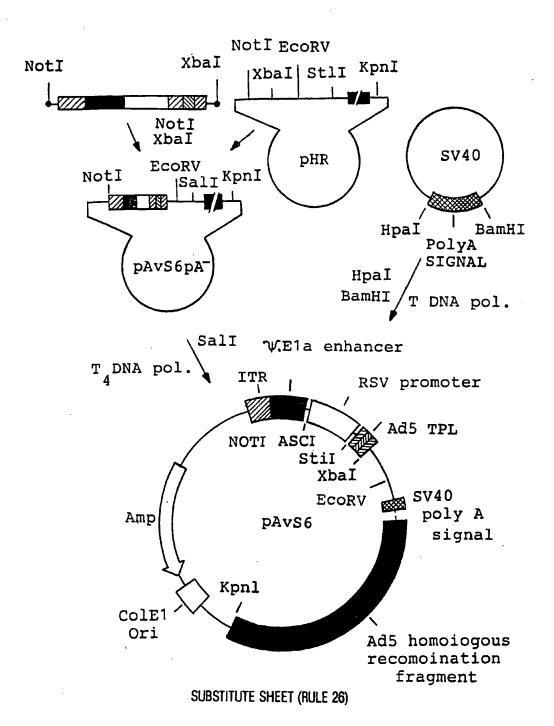
16. A process for treating lung surfactant protein deficiency states, comprising:

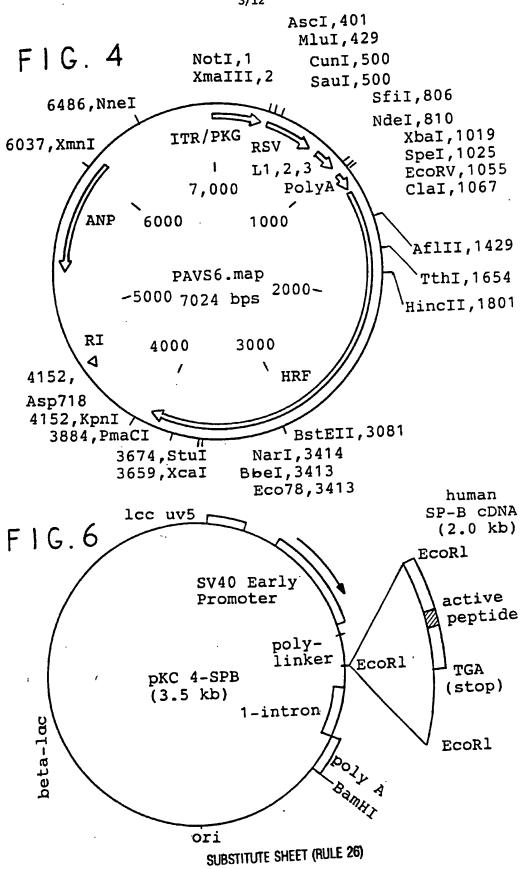
administering to a host an effective amount of the infectious viral particles of Claim 13.

- 17. The process of Claim 15 wherein said infectious viral particles are administered in an amount up to about 1013pfu.
- 18. The process of Claim 17 wherein said infectious viral particles are administered in an amount of from about 10^{7} pfu to about 10^{12} pfu.
- 19. The process of Claim 16 wherein said infectious viral particles are administered in an amount of from 1 pfu to about 10^{12} pfu.
- 20. The process of Claim 19 wherein said infectious viral particles are administered in an amount of from about $10^7 \rm pfu$ to about $10^{12} \rm pfu$.
- 21. The process of Claim 15 wherein said lung surfactant protein deficiency state is infant respiratory distress syndrome.
- The process of Claim 16 wherein said lung surfactant protein deficiency state is infant respiratory distress syndrome.
- 23. The process of Claim 15 wherein said lung surfactant protein deficiency state is adult respiratory distress syndrome.
- 24. The process of Claim 16 wherein said lung surfactant protein deficiency state is adult respiratory distress syndrome.



F1G.3





MATCH WITH FIG. 5B

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		SUE	STITUTE !	SHEET (RL	JLE 26)			•
-	61	121	181	241	301	361	421	481
AATTCCGTCA TTAAGGCAGT	GCTGCTGCCC	TGCCCAGGGC ACGGGTCCCG	AGGGCATTTC TCCCGTAACG	TGAGGACATC ACTCCTGTAG	GAGGAAGTTC CTCCTTCAAG	CAACCAAGTG GTTGGTTCAC	CTCAAACGGC GAGTTTGCCG	GGAGCCAGGG
AGCTGCAGAG TCGACGTCTC	ACGCTCTGTG TGCGAGACAC	CCTGAGTTCT GGACTCAAGA	CTACAGGAAG GATGTCCTTC	GTCCACATCC CAGGTGTAGG	CTGGAGCAGG GACCTCGTCC	CTTGACGACT GAACTGCTGA	ATCTGTATGC TAGACATACG	ATGTCAGACC TACAGTCTGG
GTGCCATGGC CACGGTACCG	GCCCAGGCAC CGGGTCCGTG	GGTGCCAAAG CCACGGTTTC	TCTGGGGACA AGACCCCTGT	TTAACAAGAT AATTGTTCTA	AGTGCAACGT TCATGTTGCA	ACTTCCCCCT TGAAGGGGGA	ACCTGGGCCT TGGACCCGGA	CCCTGCCCAA GGGACGGGTT
TGAGTCACAC ACTCAGTGTG	TGCTGCCTGG ACGACGGACC	CCTGGAGCAA GGACCTCGTT	TGTGGGAGCC	GGCCAAGGAG CCGGTTCCTC	CCTCCCCTTG	GGTCATCGAC CCAGTAGCTG	GTGCAAATCC CACGTTTAGG	ACCTCTGCGG TGGAGACGCC
CTGCTGCAGT GACGACGTCA	ACCACCTCAT TGGTGGAGTA	GCATTGCAGT CGTAACGTCA	GATGACCTAT CTACTGGATA	GCCATTTTCC CGGTAAAAGG	AAGCTGCTCA TTCGACGAGT	TACTTCCAGA ATGAAGGTCT	CGGCAGCCAG GCCGTCGGTC	GACCCTCTGC CTGGGAGACG
GGCTGCTGCT	CCTTGGCCTG	GCAGAGCCCT	GCCAAGAGTG CGGTTCTCAC	AGGACACGAT TCCTGTGCTA	TGCCCCAGTG ACGGGGTCAC	ACCAGACTGA TGGTCTGACT	AGCCAGAGCA TCGGTCTCGT	CAGACCCTCT GTCTGGGAGA

TICGITAAAC ACCICGICGI GIGCGGGTC GACGACIGGG ACCACGGGTC

F16. 5C

MATCH WITH

CCTTTTCACG

1021

GGAAAAGTGC AAGCAATTTG TGGAGCAGCA CACGCCCCAG CTGCTGACCC TGGTGCCCAG CAGCGAGCAG GCCATACTAC AGGCAATGCT CCAGGCCTGT GTTGGCTCCT GGCTGGACAG CGACGGCGCT CTGAGACTCA CGGTGGAGAC GTACAGGCAC TGGTGGGTCC GGCCCTTGTC CCGGGAACAG GICCICITAC CTACTCCGTC ATCCTGCTCG ACACGCTGCT GGGCCGCATG CTGCCCCAGC TGGTCTGCCG GATGAGGCAG TAGGACGAGC TGTGCGACGA CCCGGCGTAC GACGGGGTCG ACCAGACGCC CAGGAGAATG GGTCCACACG GCGCACCATG GAGACCACTG CCCGCCGTAG ACGGTCACGG ACCGACTCGC CCAGGTGTGC CGCGTGGTAC CTCTGGTGGC GGGCGGCATC TGCCAGTGCC TGGCTGAGCG GGCTCTGATC AAGCGGATCC AAGCCATGAT TCCCAAGGGT GCGCTACGTG TGGCAGTGGC CCGAGACTAG TTCGCTAGG TTCGGTACTA AGGGTTCCCA CGCGATGCAC ACCGTCACCG CACACAGGAT CTCTCCGAGC AGCAATICCC CATICCTCTC CCCTATIGCT GGCTCTGCAG GTGTCCTA GAGAGCTCG TCGTTÄAGGG GTAAGGAGAG GGGATAACGA CCGAGACGTC CGACCTGTTC GAGCAGGAGG GACACGACGG GCCCCGGGAG GTCCGCTCCG GACCCGGAGT CAGGCGAGGC CTGGGCCTCA GACTCTGAGT GCCACCTCTG CATGTCCGTG ACCACCCAGG GTCGCTCGTC CGGTATGATG TCCGTTACGA GGTCCGGACA ACCCCGAGGA CGGTGCTCCA TGGATGACAG CGCTGGCCCA AGGTCGCCGA GCCACGAGGT ACCTACTGTC GCGACCGGGT TCCAGCGGCT GCTGGACAAG CTCGTCCTCC CTGTGCTGCC CGGGGCCCTC MATCH WITH FIG. 5A CCTCGTCCTC GGAGCAGGAG 961 901 841 781 721 661 601 541 SUBSTITUTE SHEET (RULE 26)

AATTCAAAAT TTAAGTTTTA

GGGGTTGAAG CTGAGCCCCG

TGGAGGGTCC

GTCGTAGACC TCCTCGAGAC GTCGGAGGTG TGGATGGTGC

GGAAAAACAA GCCACTGCTT TACAGGACAG CCTTTTTGGT CGGTGACGAA ATGTCCTGTC

CGACCCGAGT

GCTGGGCTCA

1501

1561

TGTCGCTCCC

GACTCGGGGC

CCCCAACTTC

CAGCCTCCAC ACCTACCACG ACCTCCCAGG CGGCCTTGTC GGTGTGAGGC CCACACTCCG CTCACCGCCC GAGTGGCGGG GCCGGAACAG GCTCCCTGGC GTGCAAAGGA CGAGGGACCG TGTGGGACCA TGTCCAGCCC ACACCCTGGT ACAGGTCGGG CACGTTTCCT FIG. 50 GTAGGAGAGA AGCTGTCCAG TCGACAGGTC GAGAAGGGGA GTGGCGGGAG AGCCCTGTGT TCGGGACACA CATCCTCTCT CTCTPCCCCT GACCCCCTC GTTGGTGTCA GACTGTGGTC TCGGGTGAAG CTGGTGGACC ACGGGCCCCC TGCCCGGGGG CTGGGACCAT GGTGACCAGG AGCCCACTTC ATGAGAACTC TACTCTTGAG CCACTGGTCC TCAGCTCCCA GGAGCGGTCG ACGGTCCGAC TTTTCTTCGG AGTCGAGGGT CCTUGGGGT GGAGCCCCA CAGCATCTGG AGGAGCTCTG GGACGGTCCG GGCTGGAAAC GACCCTGGTA TGCCAGGCTG AAAAGAAGCC GACCACCTGG CTGACACCAG CCGACCTTTG CCTGCCAGGC FIG. SB GAGACGGGCT GTCACTTCCA TAGGTGTCGG CTCTCCCCGA CAGTGAAGGT CAACCACAGT ATCCACAGCC GGGCTGGGAT GCCCACACCA CCCGACCCTA CGGGT TGGT MATCH WITH ACAGCGAGGG TCTCCAGTGT TTCCTCGGCA CCTCGCCAGC ACAGAGTCGA AGAGGTCACA AAAGCCAAGT TTTCGGTTCA AAGGAGCCGT TGTCTCAGCT 1441 1321 1381 1081 1201 1261 1141 SUBSTITUTE SHEET (RULE 26)

CCTCACACCC ACCCCCATGC ACTCAAAGAT TGGATTTTAC AGCTACTTGC GGAGTGTGGG TGGGGGTACG TGAGTTTCTA ACCTAAAATG TCGATGAACG FIG. 5D MATCH WITH

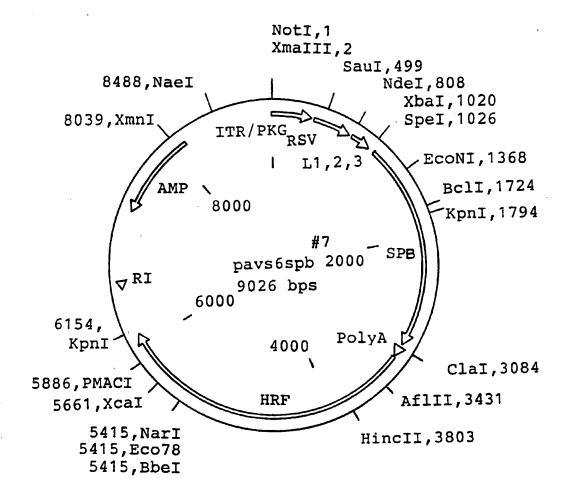
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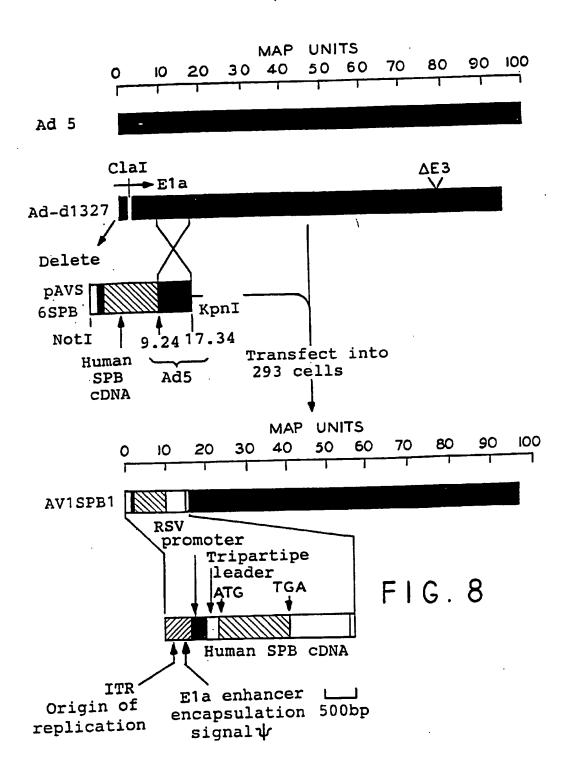
AATGCTAATA AAGTCCTATT CTCTTTTATC GGAATT TTACGATTAT TTCAGGATAA GAGAAAATAG CCTTAA

MATCH WITH FIG. SC

AAATTGTTA	CTTTAACAAT	CATGCACGCG GTACGTGCGC	TGCTGCTTGA ACGACGAACT	TAAGGTGGAC ATTCCACCTG	TATCTGCTGA ATAGACGACT	TGTCTGCACC ACAGACGTGG
ATAGACATCA GAAATTGTTA		CAAGGGTGGA CGTTCCCACCT C	CGAGATCGTC GCTC GCTCTAGCAG	TGTCTTTCTG ACAGAAAGAC	TGCTGTGATT ACGACACTAA	tgtaatacaa Acattatgtt
ACTCTAAAAG	TGAGATTTTC TATCTGTAGT	CCAGCGTAGT	CTTTCTTCCT GAAAGAAGGA	GGTTTCTTTT CCAAAGAAAA	ATTCTATTTC TAAGATAAAG	CCCATTCCTA GGGTAAGGAT
AACATACAGA ACTCTAAAAG	TTGTATGTCT	TCAGCAATTC AGTCGTTAAG	ACCGGGCAAG TGGCCCGTTC	TATAAAAAGG ATATTTTCC	TCCTAGGGTG AGGATCCCAC	AAGCTAGGGA TTCGATCCCT
AAAAAATGGG		TTTCAAAAA AAAGTTTTT	TGGGATGGCG ACCCTACCGC	TTTGTTAAGA AAACAATTCT	TGATTGAAAG ACTAACTTTC	GGGGTTGTGC CCCCAACACG
TCAGAAGAAT		AGTTAAGCTT TCAATTCGAA	TCTGGCATGA AGACCGTACT	GAGCTATTGC	TTCCAGCTTT AAGGTCGAAA	AAGTCTAGCT TTCGAGTCGA
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FIG. 7





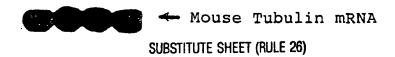
SUBSTITUTE SHEET (RULE 26)

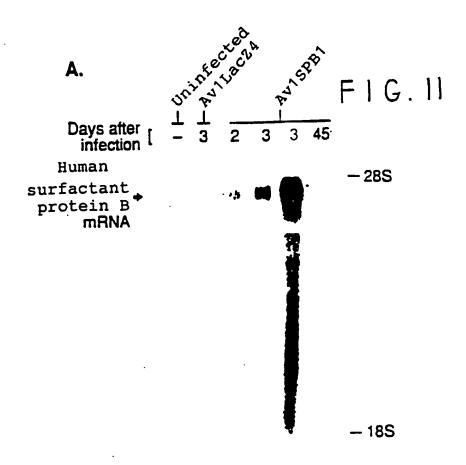
Uninfected Infected FIG. 9 Reduced Nonreduced Mr (kDa) 200 97 69 46 42 kDa ProSPB 31 - 21.5-18 kDa SPB Dimer 14.5 6-8 kDa Mature SPB 6.5 __ 2 } 3 FIG. 10

1 2 3 4 5

Mouse SPB mRNA

Human SPB mRNA





Uninfected Av1SPB1 Av1LacZ4

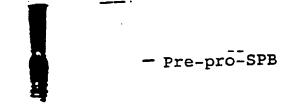


FIG.12 -

-Dimeric SPB

-- Monomeric SPB

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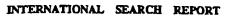
SUBSTITUTE SHEET (RULE 26)

INTERNATIONAL SEARCH REPORT

International application No. PCT/US94/03831

A. CLAS	SSIFICATION OF SUBJECT MATTER	7/00 15/00		
	A01N 63/00; A61K 37/00; C07H 17/00; C12N 5/00, Please See Extra Sheet.	•		
According to	International Patent Classification (IPC) or to both m	ational classification and IPC		
B. FIEL	DS SEARCHED	by classification symbols)		
	ocumentation searched (classification system followed 124/93A, 93R; 435/172.1, 172.3, 235.1, 240.2, 320.			
Documentati	ion searched other than minimum documentation to the	extent that such documents are included	in the fields searched	
Electronic d	ata base consulted during the international search (nam	ne of data base and, where practicable,	search terms used)	
	ee Extra Sheet.		·	
C. DOC	UMENTS CONSIDERED TO BE RELEVANT			
Category*	Citation of document, with indication, where app	propriate, of the relevant passages	Relevant to claim No.	
Y	W. Doerfler, "Adenovirus DNA", published 1986, by Martinus Nijhoff Publishing (Boston), pages 53-95, see the entire document.		1-24	
Y	Cell, Volume 68, issued 10 January al, "In Vivo Transfer of the Transmembrane Conductance Regu Epithelium", pages 143-155, see t	1-24		
Y.	Science, Volume 252, issued 19 Apet al, "Adenovirus-Mediated Trans Antitrypsin Gene to the Lung Epithe 434, see the entire document.	fer of a Recombinant α1-	1-24	
X Furt	her documents are listed in the continuation of Box C.	. See patent family annex.		
تنا	pacial categories of cited documents:	"T later document published after the in date and not in conflict with the appli	ternational filing date or priority cation but cited to understand the	
•••	ocument defining the general state of the art which is not considered be part of puriousar relevance	principle or theory underlying the m	Asimon	
.E. 01	urlier document published on or after the international filing date	"X" document of particular relevance; is considered novel or cannot be considered when the document is taken alone	ne canamed sevention cannot be lered to involve an investive step	
	nowment which may throw doubts on priority claim(s) or which is ted to catabilish the publication date of another citation or other	4	the claimed invention cannot be	
	occial reason (as specifical) occurrent referring to an oral disclosure, use, exhibition or other	considered to involve an inventor	ch documents, such combination	
	council published prior to the international filing data but inter than	being obvious to a person skilled in "A" document member of the man pates		
The Cha	s priority date classed	Date of mailing of the international a		
Date of the	actual completion of the international search	JUN 22 1994	-	
25 MAY	1994			
Name and	mailing address of the ISA/US oner of Patents and Trademarks	Authorized officer	Vyza for	
Box PCT	m, p.c. 20231	Authorized officer BRUCE CAMPELL ORGANICA BRUCE CAMPELL		
. .		Tolonhoro No. (703) 308-0196		

Facsimile No. (703) 305-3230
Form PCT/ISA/210 (second sheet)(July 1992)*



International application No. PCT/US94/03831

C (Continua	tion). DOCUMENTS CONSIDERED TO BE RELEVANT	<u> </u>	
Category*	Citation of document, with indication, where appropriate, of the relev	ant passages	Relevant to claim No
Y	DNA, Volume 8, No. 2, issued 1989, T.J. Pilot-Matias et al, "Structure and Organization of the Gene Encoding Human Pulmonary Surfactant Proteolipid SP-B", pages 75-86, see the entire document.		1-24
	Journal of Clinical Investigation, Volume 81, issued March 1988, S.D. Revak et al, "Use of Human Surfactant Low Molecular Weight Apoproteins in the Reconstitution of Surfactant Biologic Activity", pages 826-833, see the entire document.		15-24
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			•

INTERNATIONAL SEARCH REPORT

International application No. PCT/US94/03831

A. CLASSIFICATION OF SUBJECT MATTER: US CL :
424/93A, 93R; 435/172.1, 172.3, 235.1, 240.2, 320.1; 536/23.1, 23.5, 24.1, 24.2
B. FIELDS SEARCHED Electronic data bases consulted (Name of data base and where practicable terms used):
APS, BIOSIS, EXCERPTA MEDICA, DERWENT BIOTECHNOLOGY ABSTRACTS, CHEMICAL ABSTRACTS search terms: surfactant and (pulmonary or lung), adenovirus vector?, gene therapy
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